

Resource Summary Report

Generated by [kravitz2](#) on Aug 1, 2026

MetaLocGramN

RRID:SCR_003154

Type: Tool

Proper Citation

MetaLocGramN (RRID:SCR_003154)

Resource Information

URL: <http://iimcb.genesilico.pl/MetaLocGramN/>

Proper Citation: MetaLocGramN (RRID:SCR_003154)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on January 5, 2023. A tool for subcellular localization prediction of Gram-negative proteins. You can also use MetaGramLocN via SOAP. SOAP enables you to invoke our method from scripts written in your programming language of choice.

Abbreviations: MetaLocGramN

Resource Type: data analysis service, data access protocol, service resource, production service resource, software resource, analysis service resource, web service

Defining Citation: [PMID:22705560](#)

Keywords: subcellular localization, protein, prediction, sequence, analysis, gram-negative protein, gram-negative, gram-negative bacteria

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: MetaLocGramN

Resource ID: SCR_003154

Alternate IDs: OMICS_01626

Record Creation Time: 20220129T080217+0000

Record Last Update: 20260801T042530+0000

Ratings and Alerts

No rating or validation information has been found for MetaLocGramN.

No alerts have been found for MetaLocGramN.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Stuart RK, et al. (2016) Light Regimes Shape Utilization of Extracellular Organic C and N in a Cyanobacterial Biofilm. mBio, 7(3).

Moum??ne A, et al. (2015) Proteomic profiling of the outer membrane fraction of the obligate intracellular bacterial pathogen Ehrlichia ruminantium. PloS one, 10(2), e0116758.

Rubiano-Labrador C, et al. (2015) Salt Stress Induced Changes in the Exoproteome of the Halotolerant Bacterium Tistlia consotensis Deciphered by Proteogenomics. PloS one, 10(8), e0135065.